

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/500,601B
Source: PC/10
Date Processed by STIC: 9/2/05

ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 09/02/2005

PATENT APPLICATION: US/10/500,601B

TIME: 15:20:46

Input Set : A:\Substitute Sequence Listing.txt

Output Set: N:\CRF4\09012005\J500601B.raw

3 <110> APPLICANT: Repp, Reinald
 5 <120> TITLE OF INVENTION: Methods for detecting nucleic acids using solid phase bound
 6 primers and cleaving the products of a cyclic amplification
 7 reaction (Trap Release Primer Amplification (TRamp))
 9 <130> FILE REFERENCE: 221801-1010
 11 <140> CURRENT APPLICATION NUMBER: US 10/500,601B
 12 <141> CURRENT FILING DATE: 2004-06-30
 14 <150> PRIOR APPLICATION NUMBER: PCT/DE02/04754
 15 <151> PRIOR FILING DATE: 2002-12-31
 17 <150> PRIOR APPLICATION NUMBER: DE 10164219.9
 18 <151> PRIOR FILING DATE: 2001-12-31
 20 <160> NUMBER OF SEQ ID NOS: 1
 22 <170> SOFTWARE: PatentIn version 3.3
 24 <210> SEQ ID NO: 1
 25 <211> LENGTH: 32
 26 <212> TYPE: DNA
 27 <213> ORGANISM: artificial sequence
 29 <220> FEATURE:
 30 <223> OTHER INFORMATION: PCR-primer
 33 <220> FEATURE:
 34 <221> NAME/KEY: misc_feature
 35 <222> LOCATION: (1)..(1)
 36 <223> OTHER INFORMATION: n = a, t, c, or g
 38 <220> FEATURE:
 39 <221> NAME/KEY: misc_feature
 40 <222> LOCATION: (32)..(32)
 41 <223> OTHER INFORMATION: n = a, t, c, or g
 43 <400> SEQUENCE: 1

W--> 44 ntgtgtctcg agtggttttca ctgagactt tn

32

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 1,32

VERIFICATION SUMMARY

DATE: 09/02/2005

PATENT APPLICATION: US/10/500,601B

TIME: 15:20:47

Input Set : A:\Substitute Sequence Listing.txt

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L:44 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0